# RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
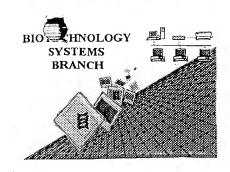BRANCH

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: *09/778,900*

Source: *1645*

Date Processed by STIC: *5-7-01*

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

## Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.
Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).
Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:
### http://www.uspto.gov/web/offices/pac/checker

1645

```
        RAW SEQUENCE LISTING              DATE: 05/07/2001
        PATENT APPLICATION:  US/09/778,900    TIME: 18:00:15

        Input Set : A:\09778900.txt
        Output Set: N:\CRF3\05072001\I778900.raw
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 5 <120> TITLE OF INVENTION: DIAGNOSTIC METHOD
 7 <130> FILE REFERENCE: 009901/0277123 - LDSG/PHM70675/US
 9 <140> CURRENT APPLICATION NUMBER: US 09/778,900
10 <141> CURRENT FILING DATE: 2001-02-08
12 <150> PRIOR APPLICATION NUMBER: GB 0004232.5
13 <151> PRIOR FILING DATE: 2000-02-24
15 <160> NUMBER OF SEQ ID NOS: 24
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22 <213> ORGANISM: Human
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25 <221> NAME/KEY: UNSURE
26 <222> LOCATION: 396
27 <223> OTHER INFORMATION: any
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32 ttagtcattg tttggaatga cttttataaaa atgctttgca ttttttagca agaccatcat 180
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34 ccaagatagg atggtagagc tcacattaca gaaatgtagt gtgggaatag taaggagtcg 300
35 tttaatagaa attgcacacc taagtgtgat gagtgtatgt gaatgtggag aagtactttc 360
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39 ctttggatga gcagtgtgag cggctccctt atgatgccag caagtgggag tttgcccggg 600
40 agagacttaa actgggtaag atatttgttc aacagattca taaacctata ctgagcacat 660
41 attacatgaa aaacactgtg ctttgagaga tgcgaaagta aactagacct gggattctac 720
42 cctccagctg ctcacagact agcaagggag atggacacaa aagtaaataa ttccaatgca 780
43 atgctcagat aacagtacaa ggtgacacgc agcacctgtt tgttcttgca acagttatta 840
44 ggcaccttct ctgagcagca gacactggtc taagccctgg agacacaaag gtgcttgcat 900
45 ctcttccctc aaagggctca gtctggagat aggtgcaaaa gtggtaagtg aagggggcg 960
46 gagagagagg cattacaagt acacgcacgc ttcataatga aactgttgag ggattagaaa 1020
47 tatgtgatcc agaacataat tgagggtggc aaggaacagt gaaatcaaca ttc      1073
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52 <212> TYPE: DNA
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61 <221> NAME/KEY: UNSURE
62 <222> LOCATION: 359
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    74 atacccacca ctcttaattt ttttttttta tctgatctaa gatgctgtct agaagcccag 240
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    86 ttaaaaatga actctgatct acttgttggt ttgttttatg ttttgctaac attgttccaa 960
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    93 ctggggaaaa ctcacaggtt tcagactaga caaacctggc atcggctctc cacagtatcc 1380
    94 tctggcatat tttcaaatct ggcccaaatc tcagaagaca tgacttcata ggagagctac 1440
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   106 <223> OTHER INFORMATION: any
   108 <400> SEQUENCE: 3
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   110 ccttcatggg actcagtata gcactctagc tccctctttt agcgtaacac tgcatactat 120
   111 ggtgttctct atgttaggaa accagagctg ctctcggaaa tgatttatag gccgtatgtt 180
   112 atctgggagg tgaccccatg gacactcggg ttgaatgtgc tttgttttca tgcccttctg 240
   113 ctcaaggccc ccttgccctc ttctagactc gacttcctct gaaatggatg gctcctgaat 300
   114 ctatctttga caaatctac agcaccaaga gcgacgtgtg tcttacgga gtattgctgt 360
   115 gggaaatctt ctccttaggt aaatttggga gaaggaagaa atcaaacagc ccagaaataa 420
   116 atgtctgcat cttctgctga atgtcctttg gttggacagc ctttagatta gaacctactg 480
   117 taacaaaaaa ctcttaaagt gtaatgggcc catgtagact ctcagatgag taatggcgta 540
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118 cgcatgtctg ccctctactg taaaagggct ttatatgatc atgaacaagg tcagaacaag 600
119 gtcatgtaaa agggctttat acgatcatga acaagggtat aaagtctgaa gcaaagtact 660
120 ttttctgtac tttgccaatt ctgcctttttc aattcctcaa cacccacacc tctaatgccc 720
121 ttaccg                                                         726
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125 <211> LENGTH: 1352
126 <212> TYPE: DNA
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137 <223> OTHER INFORMATION: any
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144 gatcactgct ctccaactat ttctgttgct gttttcccca ccagctatca gatcatgctg 300
145 gactgctggc acagagaccc aaaagaaagg ccaagatttg cagaacttgt ggaaaaacta 360
146 ggtgatttgc ttcaagcaaa tgtacaacag gtaaaactaa atttatctac atcaaaatgc 420
147 ctttgaatgt acgtcagggg ggcattttat ttgttttttt tttaagagct attaatataa 480
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157 gtttgttacg gattttttctt tttgccaaac catttaaaag gaggcttaaa gcataatagc 1080
158 actttactcc taaatacttt agtatacatt ttgtaagaag gctattgttg ctgggcacag 1140
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160 aggagttcaa aatctgcctc ggcaacatag agagacctca tcttactaaa aatttaaaaa 1260
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173 atacaagcct ggcactagca ctcgattatg ccattaaata atatttagcc gtgtagccat 180
174 gccaggtcac tttgccacct cacatccttt tcagagcacc tgataaagtc ataccacttc 240
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*(handwritten annotation:)* } position 132 is "t, not n".

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175 cctgcacatc atttctctcc tgtgccattg ggcactcaga cgagatgatg cctccagtct 300
176 ctcctacgtc tggcattctc tgatttcaca acggaccaga gtaggtccct ctgggagttt 360
177 cctcaaccct acagaatgtg aattgacaac cacgggaggc agtggcaatg ctgtcaggat 420
178 tcccaggggt cacggcgggg agatcggggc ctcaggagtt aggtgattcc tgttggtgtg 480
179 ttggttcatc ttagctggga tatggtgcct gtggtctcct gactcattag agctggatgc 540
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181 catgagcctg gaaagaatca aaacctttga agaactttta ccgaatgcca cctccatgtt 660
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183 cttccaaaca ccaacgccag taatgtagta gttcttggtc agtgagggtc tggattcagg 780
184 agtggctgaa atgacagtgt ggggaggact gacaactaga cctagctgtg cagaactaat 840
185 ttgaaagtag agttccatgc actcactcca ggacccaagt ccctgcgtgg taggaattta 900
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190 ctaaccagta tagcctggca aacaagaggt cttccagctc ttctctctaa agccctgtga 1200
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197 <213> ORGANISM: Human
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287 <213> ORGANISM: Human
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311 <210> SEQ ID NO: 19
312 <211> LENGTH: 25
313 <212> TYPE: DNA
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VERIFICATION SUMMARY
PATENT APPLICATION:  US/09/778,900

DATE: 05/07/2001
TIME: 18:00:17

Input Set : A:\09778900.txt
Output Set: N:\CRF3\05072001\I778900.raw

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L:36  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:73  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:76  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:77  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:109 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
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